

CRF Processing Date: 10/30/2001
Edited by: AN
Verified by: (STIC staff)E.O. Flare
Serial Number: 09/663,600A Changed a file from non-ASCII to ASCII**ENTERED** Changed the margins in cases where the sequence text was 'wrapped' down to the next line.

#8

EM

 Edited a formal error in the Current Application Data section, specifically: Edited the Current Application Data section with the actual current number. The number inputted by the applicant was the prior application data; or other _____ Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: _____ Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of lines; secretary initials/filename at end of file;
 page numbers throughout text; other invalid text, such as _____ Inserted mandatory headings, specifically: _____ Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____ Other:

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/663,600A

DATE: 10/30/2001
TIME: 12:34:59

Input Set : N:\jumbos\663600a.txt
Output Set: N:\CRF3\10302001\I663600A.raw

4 <110> APPLICANT: Dumas Milne Edwards, Jean-Baptiste
 5 Duclert, Aymeric
 6 Bougueleret, Lydie
 8 <120> TITLE OF INVENTION: EXTENDED CDNAS FOR SECRETED PROTEINS
 10 <130> FILE REFERENCE: 31.US3.CIP
 12 <140> CURRENT APPLICATION NUMBER: 09/663,600A
 13 <141> CURRENT FILING DATE: 2000-09-15
 15 <150> PRIOR APPLICATION NUMBER: 09/191,997
 16 <151> PRIOR FILING DATE: 1998-11-13
 18 <150> PRIOR APPLICATION NUMBER: 60/066,677
 19 <151> PRIOR FILING DATE: 1997-11-13
 21 <150> PRIOR APPLICATION NUMBER: 60/069,957
 22 <151> PRIOR FILING DATE: 1997-12-17
 24 <150> PRIOR APPLICATION NUMBER: 60/074,121
 25 <151> PRIOR FILING DATE: 1998-02-09
 27 <150> PRIOR APPLICATION NUMBER: 60/081,563
 28 <151> PRIOR FILING DATE: 1998-04-13
 30 <150> PRIOR APPLICATION NUMBER: 60/096,116
 31 <151> PRIOR FILING DATE: 1998-08-10
 33 <150> PRIOR APPLICATION NUMBER: 60/099,273
 34 <151> PRIOR FILING DATE: 1998-09-04
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 56 <213> ORGANISM: Artificial Sequence
 W--> 57 <220> FEATURE:
 58 <223> OTHER INFORMATION: in vitro transcription product
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 64 <212> TYPE: DNA
 65 <213> ORGANISM: Artificial Sequence

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/663,600A
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 Input Set : N:\jumbos\663600a.txt
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  90 <211> LENGTH: 25  

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  92 <213> ORGANISM: Artificial Sequence  

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  94 <223> OTHER INFORMATION: Alpha globin gene primer GLO-As  

W--> 95 <400> SEQUENCE: 6  

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  100 <212> TYPE: DNA  

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  103 <223> OTHER INFORMATION: Dehydrogenase gene primer 3 DH-S  

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  105 agtgattccct gctactttgg atggc 25  

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  109 <212> TYPE: DNA  

  110 <213> ORGANISM: Artificial Sequence  

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  117 <211> LENGTH: 25  

  118 <212> TYPE: DNA  

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RAW SEQUENCE LISTING DATE: 10/30/2001
 PATENT APPLICATION: US/09/663,600A TIME: 12:34:59

Input Set : N:\jumbos\663600a.txt
 Output Set: N:\CRF3\10302001\I663600A.raw

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134 <210> SEQ ID NO: 11
135 <211> LENGTH: 25
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168 <221> NAME/KEY: misc_feature
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170 <223> OTHER INFORMATION: n=a, g, c or t
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173 tttttvnn 67
175 <210> SEQ ID NO: 15
  
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RAW SEQUENCE LISTING DATE: 10/30/2001
 PATENT APPLICATION: US/09/663,600A TIME: 12:34:59

Input Set : N:\jumbos\663600a.txt
 Output Set: N:\CRF3\10302001\I663600A.raw

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186 <212> TYPE: DNA
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207 <222> LOCATION: complement(110..145)
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211 <222> LOCATION: complement(196..229)
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214 <221> NAME/KEY: sig_peptide
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216 <223> OTHER INFORMATION: Von Heijne matrix
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218 <221> NAME/KEY: misc_feature
219 <222> LOCATION: 290
220 <223> OTHER INFORMATION: n=a, g, c or t
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222 aatatrarac agtataacaata ttccaggccc artcaattgc catttctcat aacagcgta 60
223 gagagaaaaga actgactgar acgtttgag atg aag aaa gtt ctc ctc ctg atc 113
224 Met Lys Lys Val Leu Leu Ile
225 -15 -10
226 aca gcc atc ttg gca gtg gct gtw ggt ttc cca gtc tct caa gac cag 161
  
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/663,600A

DATE: 10/30/2001
TIME: 12:35:00

Input Set : N:\jumbos\663600a.txt
Output Set: N:\CRF3\10302001\I663600A.raw

227	Thr Ala Ile Leu Ala Val Ala Val Gly Phe Pro Val Ser Gln Asp Gln			
228	-5	1	5	
229	gaa cga gaa aaa aga agt atc agt gac agc gat gaa tta gct tca ggr	209		
230	Glu Arg Glu Lys Arg Ser Ile Ser Asp Ser Asp Glu Leu Ala Ser Gly			
231	10	15	20	
232	wtt ttt gtg ttc cct tac cca tat cca ttt cgc cca ctt cca cca att	257		
W--> 233	Xaa Phe Val Phe Pro Tyr Pro Tyr Pro Phe Arg Pro Leu Pro Pro Ile			
234	25	30	35	
W--> 235	cca ttt cca aga ttt cca tgg ttt aga cgt aan ttt cct att cca ata	305		
W--> 236	Pro Phe Pro Arg Phe Pro Trp Phe Arg Arg Xaa Phe Pro Ile Pro Ile			
237	40	45	50	55
238	cct gaa tct gcc cct aca act ccc ctt cct agc gaa aag taaacaaraaa	354		
239	Pro Glu Ser Ala Pro Thr Thr Pro Leu Pro Ser Glu Lys			
240	60	65		
241	ggaaaaagtca crataaacct ggtcacctga aattgaaatt gagccacttc cttgaaraat	414		
242	caaaaattcct gttataaaaaa raaaaacaaa tgtaattgaa atagcacaca gcattctcta	474		
243	gtcaatatct ttagtgatct tcttaataaa acatgaaagc aaaaaaaaaaa aa	526		
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248	<213> ORGANISM: Homo Sapiens			
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251	<222> LOCATION: 1..17			
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253	score 8.2			
254	seq LLLITAILAVAVG/FP			
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258	Gly			
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274	<222> LOCATION: 56..113			
275	<223> OTHER INFORMATION: blastn			
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277	<221> NAME/KEY: misc_feature			

PNJ Use of n and/or Xaa has been detected in the Sequence Listing.
Review the Sequence Listing to insure a corresponding explanation is presented in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/663,600A

DATE: 10/30/2001
TIME: 12:35:01

Input Set : N:\jumbos\663600a.txt
Output Set: N:\CRF3\10302001\I663600A.raw

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L:104 M:283 W: Missing Blank Line separator, <400> field identifier
L:111 M:283 W: Missing Blank Line separator, <220> field identifier
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VERIFICATION SUMMARY
PATENT APPLICATION: US/09/663,600A

DATE: 10/30/2001
TIME: 12:35:01

Input Set : N:\jumbos\663600a.txt
Output Set: N:\CRF3\10302001\I663600A.raw

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/663,600A

DATE: 10/30/2001

TIME: 12:35:01

Input Set : N:\jumbos\663600a.txt

Output Set: N:\CRF3\10302001\I663600A.raw

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1600

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/663,600A

DATE: 10/17/2001
TIME: 12:26:46

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Output Set: N:\CRF3\10172001\I663600A.raw

4 <110> APPLICANT: Dumas Milne Edwards, Jean-Baptiste
 5 Duclert, Aymeric
 6 Bougueret, Lydie
 8 <120> TITLE OF INVENTION: EXTENDED CDNAS FOR SECRETED PROTEINS P.2
 10 <130> FILE REFERENCE: 31.US3.CIP
 12 <140> CURRENT APPLICATION NUMBER: 09/663,600A
 13 <141> CURRENT FILING DATE: 2000-09-15
 15 <150> PRIOR APPLICATION NUMBER: 09/191,997
 16 <151> PRIOR FILING DATE: 1998-11-13
 18 <150> PRIOR APPLICATION NUMBER: 60/066,677
 19 <151> PRIOR FILING DATE: 1997-11-13
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 28 <151> PRIOR FILING DATE: 1998-04-13
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 33 <150> PRIOR APPLICATION NUMBER: 60/099,273
 34 <151> PRIOR FILING DATE: 1998-09-04
 36 <160> NUMBER OF SEQ ID NOS: 229
 38 <170> SOFTWARE: Patent.pm

Does Not Comply
Corrected Diskette Needed

Delete End of file
New ASCII Text

ERRORED SEQUENCES

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 12244 Lys Leu Arg Lys Phe Arg Phe Arg Lys Glu Thr Asp Asn Ala Ala Ile
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 12252 Arg Val Ser Tyr Pro Leu Cys Phe Ile Phe Ser Ser Pro Val Gly Cys
 12253 85 90 95
 12254 Lys Pro Glu Gln Gln Met Met Tyr Ala Gly Ser Lys Asn Arg Leu Val
 12255 100 105 110
 12256 Gln Thr Ala Glu Leu Thr Lys Val Phe Glu Ile Arg Thr Thr Asp Asp

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/663,600A

DATE: 10/17/2001
TIME: 12:26:49

Input Set : D:\Seqlist.txt
Output Set: N:\CRF3\10172001\I663600A.raw

12257 115 120 125
12258 Leu Thr Glu Ala Trp Leu Gln Glu Lys Leu Ser Phe Phe Arg
12259 130 135 140

E--> 12260 1

Errored Delete End of file Non ASCII Text

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/663,600A

DATE: 10/17/2001
TIME: 12:26:50

Input Set : D:\Seqlist.txt
Output Set: N:\CRF3\10172001\I663600A.raw

L:44 M:283 W: Missing Blank Line separator, <220> field identifier
L:46 M:283 W: Missing Blank Line separator, <220> field identifier
L:50 M:283 W: Missing Blank Line separator, <400> field identifier
L:57 M:283 W: Missing Blank Line separator, <220> field identifier
L:59 M:283 W: Missing Blank Line separator, <400> field identifier
L:66 M:283 W: Missing Blank Line separator, <220> field identifier
L:68 M:283 W: Missing Blank Line separator, <400> field identifier
L:75 M:283 W: Missing Blank Line separator, <220> field identifier
L:77 M:283 W: Missing Blank Line separator, <400> field identifier
L:84 M:283 W: Missing Blank Line separator, <220> field identifier
L:86 M:283 W: Missing Blank Line separator, <400> field identifier
L:93 M:283 W: Missing Blank Line separator, <220> field identifier
L:95 M:283 W: Missing Blank Line separator, <400> field identifier
L:102 M:283 W: Missing Blank Line separator, <220> field identifier
L:104 M:283 W: Missing Blank Line separator, <400> field identifier
L:111 M:283 W: Missing Blank Line separator, <220> field identifier
L:113 M:283 W: Missing Blank Line separator, <400> field identifier
L:120 M:283 W: Missing Blank Line separator, <220> field identifier
L:122 M:283 W: Missing Blank Line separator, <400> field identifier
L:129 M:283 W: Missing Blank Line separator, <220> field identifier
L:131 M:283 W: Missing Blank Line separator, <400> field identifier
L:138 M:283 W: Missing Blank Line separator, <220> field identifier
L:140 M:283 W: Missing Blank Line separator, <400> field identifier
L:147 M:283 W: Missing Blank Line separator, <220> field identifier
L:149 M:283 W: Missing Blank Line separator, <400> field identifier
L:156 M:283 W: Missing Blank Line separator, <220> field identifier
L:158 M:283 W: Missing Blank Line separator, <400> field identifier
L:165 M:283 W: Missing Blank Line separator, <220> field identifier
L:167 M:283 W: Missing Blank Line separator, <220> field identifier
L:171 M:283 W: Missing Blank Line separator, <400> field identifier
L:173 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:179 M:283 W: Missing Blank Line separator, <220> field identifier
L:181 M:283 W: Missing Blank Line separator, <400> field identifier
L:188 M:283 W: Missing Blank Line separator, <220> field identifier
L:190 M:283 W: Missing Blank Line separator, <400> field identifier
L:197 M:283 W: Missing Blank Line separator, <220> field identifier
L:201 M:283 W: Missing Blank Line separator, <220> field identifier
L:205 M:283 W: Missing Blank Line separator, <220> field identifier
L:209 M:283 W: Missing Blank Line separator, <220> field identifier
L:213 M:283 W: Missing Blank Line separator, <220> field identifier
L:217 M:283 W: Missing Blank Line separator, <220> field identifier
L:221 M:283 W: Missing Blank Line separator, <400> field identifier
L:233 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:235 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:236 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:249 M:283 W: Missing Blank Line separator, <220> field identifier
L:255 M:283 W: Missing Blank Line separator, <400> field identifier
L:264 M:283 W: Missing Blank Line separator, <220> field identifier

VERIFICATION SUMMARY
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Input Set : D:\Seqlist.txt
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L:268 M:283 W: Missing Blank Line separator, <220> field identifier
L:272 M:283 W: Missing Blank Line separator, <220> field identifier
L:276 M:283 W: Missing Blank Line separator, <220> field identifier
L:280 M:283 W: Missing Blank Line separator, <220> field identifier
L:284 M:283 W: Missing Blank Line separator, <220> field identifier
L:288 M:283 W: Missing Blank Line separator, <220> field identifier
L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:328 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:379 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
L:429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:435 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:441 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:463 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:500 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:503 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:506 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:568 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:571 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:623 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:31 ✓
L:810 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:34
L:927 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:928 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:964 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:37
L:1272 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40
L:1492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1494 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1495 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1498 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42
L:1804 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:1810 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45
L:2351 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48
L:2496 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49
L:2540 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49
L:2541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49
L:2542 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49
L:2600 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50
L:2926 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:2941 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:2950 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:2953 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:2956 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:52
L:3104 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:53
L:3154 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:54
L:3401 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:3403 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:3404 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56
L:3407 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:56

VERIFICATION SUMMARY
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Input Set : D:\Seqlist.txt
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L:3651 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58
L:3660 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:58
L:3838 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:3847 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:3859 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:3865 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:59
L:12260 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:229 Q/V